

**IFWO** 

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/789,956**DATE: 07/28/2004
TIME: 16:46:07

Input Set : A:\She0081.app

Output Set: N:\CRF4\07282004\J789956.raw

3 <110> APPLICANT: BOSSARD, MARY J. 4 BENTLEY, MICHAEL D. 6 <120> TITLE OF INVENTION: POLYMER-FACTOR VIII MOIETY CONJUGATES 8 <130> FILE REFERENCE: SHE0081.00 10 <140> CURRENT APPLICATION NUMBER: 10/789,956 11 <141> CURRENT FILING DATE: 2004-02-26 13 <150> PRIOR APPLICATION NUMBER: 60/450,578 14 <151> PRIOR FILING DATE: 2003-02-26 ENTERED 16 <160> NUMBER OF SEQ ID NOS: 2 18 <170> SOFTWARE: PatentIn Ver. 3.2 20 <210> SEQ ID NO: 1 21 <211> LENGTH: 2351 22 <212> TYPE: PRT 23 <213> ORGANISM: Homo sapiens 25 <400> SEQUENCE: 1 26 Met Gln Ile Glu Leu Ser Thr Cys Phe Phe Leu Cys Leu Leu Arg Phe 5 29 Cys Phe Ser Ala Thr Arg Arg Tyr Tyr Leu Gly Ala Val Glu Leu Ser 25 32 Trp Asp Tyr Met Gln Ser Asp Leu Gly Glu Leu Pro Val Asp Ala Arg 35 Phe Pro Pro Arg Val Pro Lys Ser Phe Pro Phe Asn Thr Ser Val Val 38 Tyr Lys Lys Thr Leu Phe Val Glu Phe Thr Asp His Leu Phe Asn Ile 70 41 Ala Lys Pro Arg Pro Pro Trp Met Gly Leu Leu Gly Pro Thr Ile Gln 85 90 44 Ala Glu Val Tyr Asp Thr Val Val Ile Thr Leu Lys Asn Met Ala Ser 100 105 47 His Pro Val Ser Leu His Ala Val Gly Val Ser Tyr Trp Lys Ala Ser 120 50 Glu Gly Ala Glu Tyr Asp Asp Gln Thr Ser Gln Arg Glu Lys Glu Asp 135 53 Asp Lys Val Phe Pro Gly Gly Ser His Thr Tyr Val Trp Gln Val Leu 150 155 56 Lys Glu Asn Gly Pro Met Ala Ser Asp Pro Leu Cys Leu Thr Tyr Ser 170 59 Tyr Leu Ser His Val Asp Leu Val Lys Asp Leu Asn Ser Gly Leu Ile 180 185 62 Gly Ala Leu Leu Val Cys Arg Glu Gly Ser Leu Ala Lys Glu Lys Thr 195 200 205 65 Gln Thr Leu His Lys Phe Ile Leu Leu Phe Ala Val Phe Asp Glu Gly 66 210 215 220

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68 Lys Ser Trp His Ser Glu Thr Lys Asn Ser Leu Met Gln Asp Arg Asp 235 69 225 230 71 Ala Ala Ser Ala Arg Ala Trp Pro Lys Met His Thr Val Asn Gly Tyr 245 250 74 Val Asn Arg Ser Leu Pro Gly Leu Ile Gly Cys His Arg Lys Ser Val 260 265 77 Tyr Trp His Val Ile Gly Met Gly Thr Thr Pro Glu Val His Ser Ile 280 80 Phe Leu Glu Gly His Thr Phe Leu Val Arg Asn His Arg Gln Ala Ser 295 83 Leu Glu Ile Ser Pro Ile Thr Phe Leu Thr Ala Gln Thr Leu Leu Met 310 315 86 Asp Leu Gly Gln Phe Leu Leu Phe Cys His Ile Ser Ser His Gln His 325 330 89 Asp Gly Met Glu Ala Tyr Val Lys Val Asp Ser Cys Pro Glu Glu Pro 340 345 92 Gln Leu Arg Met Lys Asn Asn Glu Glu Ala Glu Asp Tyr Asp Asp Asp 360 365 95 Leu Thr Asp Ser Glu Met Asp Val Val Arg Phe Asp Asp Asp Asn Ser 370 375 98 Pro Ser Phe Ile Gln Ile Arg Ser Val Ala Lys Lys His Pro Lys Thr 390 395 101 Trp Val His Tyr Ile Ala Ala Glu Glu Glu Asp Trp Asp Tyr Ala Pro 104 Leu Val Leu Ala Pro Asp Asp Arg Ser Tyr Lys Ser Gln Tyr Leu Asn 425 107 Asn Gly Pro Gln Arg Ile Gly Arg Lys Tyr Lys Lys Val Arg Phe Met 110 Ala Tyr Thr Asp Glu Thr Phe Lys Thr Arg Glu Ala Ile Gln His Glu 455 113 Ser Gly Ile Leu Gly Pro Leu Leu Tyr Gly Glu Val Gly Asp Thr Leu 475 470 116 Leu Ile Ile Phe Lys Asn Gln Ala Ser Arg Pro Tyr Asn Ile Tyr Pro 485 490 119 His Gly Ile Thr Asp Val Arg Pro Leu Tyr Ser Arg Arg Leu Pro Lys 500 505 122 Gly Val Lys His Leu Lys Asp Phe Pro Ile Leu Pro Gly Glu Ile Phe 520 515 125 Lys Tyr Lys Trp Thr Val Thr Val Glu Asp Gly Pro Thr Lys Ser Asp 535 540 128 Pro Arg Cys Leu Thr Arg Tyr Tyr Ser Ser Phe Val Asn Met Glu Arg 550 555 131 Asp Leu Ala Ser Gly Leu Ile Gly Pro Leu Leu Ile Cys Tyr Lys Glu 570 134 Ser Val Asp Gln Arg Gly Asn Gln Ile Met Ser Asp Lys Arg Asn Val 585 137 Ile Leu Phe Ser Val Phe Asp Glu Asn Arg Ser Trp Tyr Leu Thr Glu 600 140 Asn Ile Gln Arg Phe Leu Pro Asn Pro Ala Gly Val Gln Leu Glu Asp

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1/1		610					615					620				
	Pro		Dhe	Gln				Tle	Met	His	Ser		Δsn	Glv	Tur	Val
	625	GIU	THE	0111	HIU	630	11011	110	rice		635		11011	O <sub>T</sub>	- 7 -	640
	Phe	Acn	Ser	T. <del>-</del> 11	Gln		Ser	Val	Cvs	T.e11		Glu	Val	Δla·	Tur	
147	FIIC	Чал	DCI	шси	645	шси	DCI	Vul	Cyb	650	1115	O.L.	•	1114	655	115
	Tyr	τlΔ	T.011	Ser		Glv	Δla	Gln	Thr		Phe	T.em	Ser	Val		Phe
150	1 Y 1	110	anc a	660	110	017	1114	0.1.1	665	шь			DC_	670		
	Ser	Glv	ጥህን		Dhe	Lve	Hig	Lve		Val	Tur	Glu	Asn		Len	Thr
153	DCI	Ory	675	1111	1110	цуБ	1110	680		• • • •	-1-	014	685			
	Leu	Dha		Dho	Ser	Glv	Glu		Val	Phe	Met	Ser		Glu	Asn	Pro
156	пси	690	110	1110	DCI	OT Y	695	1111	• • • •	1110	1100	700		0_0		
	Gly		Trn	Tle	Len	Glv		His	Asn	Ser	Asp		Ara	Asn	Ara	Glv
	705	шец	115			710	O <sub>I</sub> D				715		5		5	720
	Met	Thr	Δla	Len	Leu		Val	Ser	Ser	Cvs		Lvs	Asn	Thr	Glv	
162	1100				725	270			-00-	730		-1-			735	
	Tyr	Tvr	Glu	Asp		Tvr	Glu	Asp	Ile		Ala	Tvr	Leu	Leu		Lvs
165						-1-			745			2		750		1
	Asn				Glu	Pro	Ara	Ser		Ser	Gln	Asn	Ser	Arq	His	Pro
168			755				5	760					765	,		
	Ser	Thr		Gln	Lvs	Gln	Phe	Asn	Ala	Thr	Thr	Ile	Pro	Glu	Asn	Asp
171		770			_1		775					780				-
	Ile		Lvs	Thr	asA	Pro	Trp	Phe	Ala	His	Arq	Thr	Pro	Met	Pro	Lys
	785		4		_	790	-				795					800
	Ile	Gln	Asn	Val	Ser	Ser	Ser	Asp	Leu	Leu	Met	Leu	Leu	Arg	Gln	Ser
177					805			~		810				_	815	
179	Pro	Thr	Pro	His	Gly	Leu	Ser	Leu	Ser	Asp	Leu	Gln	Glu	Ala	Lys	Tyr
180				820	_				825	_				830		
182	Glu	Thr	Phe	Ser	Asp	Asp	Pro	Ser	Pro	Gly	Ala	Ile	Asp	Ser	Asn	Asn
183			835					840					845			
185	Ser	Leu	Ser	Glu	Met	Thr	His	Phe	Arg	Pro	Gln	Leu	His	His	Ser	Gly
186		850					855					860	•			
188	Asp	Met	Val	Phe	Thr	Pro	Glu	Ser	Gly	Leu	Gln	Leu	Arg	Leu	Asn	Glu
	865					870					875					880
191	Lys	Leu	Gly	Thr	Thr	Ala	Ala	Thr	Glu	Leu	Lys	Lys	Leu	Asp	Phe	Lys
192					885					890					895	
194	Val	Ser	Ser	Thr	Ser	Asn	Asn	Leu	Ile	Ser	Thr	Ile	Pro	Ser	Asp	Asn
195				900					905					910		
197	Leu	Ala	Ala	Gly	Thr	Asp	Asn	Thr	Ser	Ser	Leu	Gly	Pro	Pro	Ser	Met
198			915					920					925			
200	Pro	Val	His	Tyr	Asp	Ser	Gln	Leu	Asp	Thr	Thr	Leu	Phe	Gly	Lys	Lys
201		930					935					940				
	Ser	Ser	Pro	Leu	Thr		Ser	Gly	Gly	Pro		Ser	Leu	Ser	Glu	
	945					950					955					960
206	Asn	Asn	Asp	Ser	Lys	Leu	Leu	Glu	Ser	Gly	Leu	Met	Asn	Ser		Glu
207					965					970					975	
	Ser	Ser	$\operatorname{Trp}$		Lys	Asn	Val	Ser		Thr	Glu	Ser	Gly		Leu	Phe
210				980					985			_		990		_ •
	Lys	Gly		Arg	Ala	His			Ala	Leu	Leu			Asp	Asn	Ala
213			995				:	1000					1005			

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215 Leu Phe Lys Val Ser Ile Ser Leu Leu Lys Thr Asn Lys Thr Ser Asn 216 1010 1015 1020 218 Asn Ser Ala Thr Asn Arg Lys Thr His Ile Asp Gly Pro Ser Leu Leu 219 1025 1030 1035 221 Ile Glu Asn Ser Pro Ser Val Trp Gln Asn Ile Leu Glu Ser Asp Thr 1045 1050 1055 224 Glu Phe Lys Lys Val Thr Pro Leu Ile His Asp Arg Met Leu Met Asp 225 1060 1065 1070 227 Lys Asn Ala Thr Ala Leu Arg Leu Asn His Met Ser Asn Lys Thr Thr 228 1075 1080 1085 230 Ser Ser Lys Asn Met Glu Met Val Gln Gln Lys Lys Glu Gly Pro Ile 1090 231 1095 1100 233 Pro Pro Asp Ala Gln Asn Pro Asp Met Ser Phe Phe Lys Met Leu Phe 1110 1115 236 Leu Pro Glu Ser Ala Arg Trp Ile Gln Arg Thr His Gly Lys Asn Ser 1130 1125 239 Leu Asn Ser Gly Gln Gly Pro Ser Pro Lys Gln Leu Val Ser Leu Gly 240 1140 1145 242 Pro Glu Lys Ser Val Glu Gly Gln Asn Phe Leu Ser Glu Lys Asn Lys 243 1155 1160 1165 245 Val Val Val Gly Lys Gly Glu Phe Thr Lys Asp Val Gly Leu Lys Glu 1170 1175 1180 248 Met Val Phe Pro Ser Ser Arg Asn Leu Phe Leu Thr Asn Leu Asp Asn 249 1185 1190 1195 1200 251 Leu His Glu Asn Asn Thr His Asn Gln Glu Lys Lys Ile Gln Glu Glu 1210 1205 254 Ile Glu Lys Lys Glu Thr Leu Ile Gln Glu Asn Val Val Leu Pro Gln 1220 1225 257 Ile His Thr Val Thr Gly Thr Lys Asn Phe Met Lys Asn Leu Phe Leu 1240 1245 258 1235 260 Leu Ser Thr Arg Gln Asn Val Glu Gly Ser Tyr Asp Gly Ala Tyr Ala 1255 1260 263 Pro Val Leu Gln Asp Phe Arg Ser Leu Asn Asp Ser Thr Asn Arg Thr 1275 1280 264 1265 1270 266 Lys Lys His Thr Ala His Phe Ser Lys Lys Gly Glu Glu Glu Asn Leu 267 1285 1290 269 Glu Gly Leu Gly Asn Gln Thr Lys Gln Ile Val Glu Lys Tyr Ala Cys 270 1300 1305 1310 272 Thr Thr Arg Ile Ser Pro Asn Thr Ser Gln Gln Asn Phe Val Thr Gln 273 1315 1320 1325 275 Arg Ser Lys Arg Ala Leu Lys Gln Phe Arg Leu Pro Leu Glu Glu Thr 1330 1335 278 Glu Leu Glu Lys Arg Ile Ile Val Asp Asp Thr Ser Thr Gln Trp Ser 1350 279 1345 1355 281 Lys Asn Met Lys His Leu Thr Pro Ser Thr Leu Thr Gln Ile Asp Tyr 1365 1370 284 Asn Glu Lys Glu Lys Gly Ala Ile Thr Gln Ser Pro Leu Ser Asp Cys 285 1380 1385 287 Leu Thr Arg Ser His Ser Ile Pro Gln Ala Asn Arg Ser Pro Leu Pro 
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288	1	1400					1405							
290	Ile Ala	Lys Va	al Ser	Ser	Phe	Pro	Ser	Ile	Arg	Pro	Ile	Tyr	Leu	Thr
291	1410			1	415				1	.420				
293	Arg Val	Leu Ph	ne Gln	Asp	Asn	Ser	Ser	His	Leu	Pro	Ala	Ala	Ser	Tyr
294	1425		:	1430				1	435				1	L <b>44</b> 0
296	Arg Lys	Lys As	sp Ser	Gly	Val	Gln	Glu	Ser	Ser	His	Phe	Leu	Gln	Gly
297			1445				1	450				1	L <b>4</b> 55	
299	Ala Lys	Lys As	n Asn	Leu	Ser	Leu	Ala	Ile	Leu	Thr	Leu	Glu	Met	Thr
300		146					465					L470		
302	Gly Asp	Gln Ar	g Glu	Val	Gly	Ser	Leu	Gly	Thr	Ser	Ala	Thr	Asn	Ser
303		475				L480					L <b>4</b> 85			
305	Val Thr	Tyr Ly	s Lys	Val	Glu	Asn	Thr	Val	Leu	Pro	Lys	Pro	Asp	Leu
306					495					L500		_		
	Pro Lys	Thr Se			Val	Glu	Leu			Lys	Val	His		
	1505			1510					1515		_	_		L520
	Gln Lys	Asp Le		Pro	Thr	GIu			Asn	GLY	Ser			Hıs
312			1525		_	_		1530	~ 7	_,	~ 7		1535	
	Leu Asp			GIY	Ser			GIn	GIY	Thr			Ата	TTe
315		154			7		1545	T	77. 7	D		L550	7	777
	Lys Trp		lu Ala	Asn		Pro 1560	GIY	_			Pne 1565	Leu	Arg	vai
318	Ala Thr	.555	r Cor	ח ד ת								7 cn	Dro	Lou
321		GIU Se	er ser		.575	TIIT	PIO	per	-	1580	пеа	дар	FIO	пец
	Ala Trp	Acn Ac	en Hie			Thr	Gln	Tle			G] 11	Glu	Trn	Lws
	1585	Asp As		1590	GIY				1595	цур	Gru	GIU		L600
	Ser Gln	Glu Ia			Glu					Lvs	Lvs	Lvs		
327		<sub>1</sub>	1605			-1-		L610		-1-	-2-		L615	
	Ile Leu	Ser Le		Ala	Cvs	Glu			His	Ala	Ile			Ile
330		162			1		L625					L630		
	Asn Glu	Gly Gl	ln Asn	Lys	Pro	Glu	Ile	Glu	Val	Thr	Trp	Ala	Lys	Gln
333		.635		_	1						L645			
335	Gly Arg	Thr G	lu Arg	Leu	Cys	Ser	Gln	Asn	Pro	Pro	Val	Leu	Lys	Arg
336	1650			1	655				1	L660				
338	His Gln	Arg Gl	lu Ile	Thr	Arg	Thr	Thr	Leu	Gln	Ser	Asp	Gln	Glu	Glu
	1665			1670					L675					L680
341	Ile Asp	Tyr As	sp Asp	Thr	Ile	Ser			Met	Lys	Lys			Phe
342			1685					L690				_	L695	
	Asp Ile	_	_	Asp	Glu			Ser	Pro	Arg			Gln	Lys
345		170	ነስ				705				-	L710		
2/7							L705							
34/	Lys Thr			Phe		Ala		Val	Glu	_	Leu		Asp	Tyr
348	1	Arg Hi .715	is Tyr		1	Ala 1720	Ala			- :	Leu L725	Trp	_	_
348 350	Gly Met	Arg Hi .715	is Tyr	Pro	His	Ala 1720	Ala		Asn	Arg	Leu L725	Trp	_	_
348 350 351	1 Gly Met 1730	Arg Hi .715 Ser Se	ls Tyr er Ser	Pro 1	His .735	Ala 1720 Val	Ala Leu	Arg	Asn	Arg L740	Leu 1725 Ala	Trp Gln	Ser	Gly
348 350 351 353	Gly Met 1730 Ser Val	Arg Hi .715 Ser Se	is Tyr er Ser in Phe	Pro 1 Lys	His .735	Ala 1720 Val	Ala Leu	Arg Phe	Asn Gln	Arg L740	Leu 1725 Ala	Trp Gln	Ser Asp	Gly Gly
348 350 351 353 354	Gly Met 1730 Ser Val 1745	Arg Hi 715 Ser Se Pro Gl	er Ser	Pro 1 Lys 1750	His 735 Lys	Ala 1720 Val Val	Ala Leu Val	Arg Phe	Asn Gln L755	Arg 1740 Glu	Leu 1725 Ala Phe	Trp Gln Thr	Ser Asp	Gly Gly L760
348 350 351 353 354 356	Gly Met 1730 Ser Val	Arg Hi 715 Ser Se Pro Gl	er Ser In Phe	Pro 1 Lys 1750	His 735 Lys	Ala 1720 Val Val	Ala Leu Val Gly	Arg Phe Glu	Asn Gln L755	Arg 1740 Glu	Leu 1725 Ala Phe	Trp Gln Thr His	Ser Asp Leu	Gly Gly L760
348 350 351 353 354 356 357	Gly Met 1730 Ser Val 1745 Ser Phe	Arg Hi 715 Ser Se Pro Gl	er Ser In Phe In Pro 1765	Pro 1 Lys 1750 Leu	His 735 Lys Tyr	Ala 1720 Val Val Arg	Ala Leu Val Gly	Arg Phe Glu 1770	Asn Gln L755 Leu	Arg 1740 Glu Asn	Leu 1725 Ala Phe Glu	Trp Gln Thr His	Ser Asp Leu L775	Gly Gly 1760 Gly
348 350 351 353 354 356 357	Gly Met 1730 Ser Val 1745	Arg Hi 715 Ser Se Pro Gl	er Ser In Phe In Pro 1765	Pro 1 Lys 1750 Leu	His 735 Lys Tyr	Ala 1720 Val Val Arg	Ala Leu Val Gly	Arg Phe Glu 1770	Asn Gln L755 Leu	Arg 1740 Glu Asn	Leu 1725 Ala Phe Glu Asn	Trp Gln Thr His	Ser Asp Leu L775	Gly Gly 1760 Gly

VERIFICATION SUMMARY

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